

WEST Search History

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DATE: Monday, December 06, 2004

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			<i>DB=PGPB,USPT,USOC,EPAB,JPAB,DWPI,TDBD; PLUR=YES; OP=ADJ</i>	
<input type="checkbox"/>	L25	L23 and L16		14
<input type="checkbox"/>	L24	L23 and L4		2
<input type="checkbox"/>	L23	((domain near5 specific) near5 (evolution or evolve or evolved or evolves or evolving or shuffle or shuffled or shuffles or shuffling))		30
<input type="checkbox"/>	L22	L13 and L4		12
<input type="checkbox"/>	L21	L19 and L4		2
<input type="checkbox"/>	L20	L19 and L2		54
<input type="checkbox"/>	L19	L17 and L3		70
<input type="checkbox"/>	L18	L17 same L3		3
<input type="checkbox"/>	L17	L16 same L1		213
<input type="checkbox"/>	L16	(homology near5 (clamp or domain or region))		19158
<input type="checkbox"/>	L15	L14 and (homology near5 (clamp or domain or region))		116
<input type="checkbox"/>	L14	L13 and L2		174
<input type="checkbox"/>	L13	L1 and L3		521
<input type="checkbox"/>	L12	L11 and L4		2
<input type="checkbox"/>	L11	L1 same L3		45
<input type="checkbox"/>	L10	L9 and L2		3
<input type="checkbox"/>	L9	L1 and L4		46
<input type="checkbox"/>	L8	L1 near50 L4		0
<input type="checkbox"/>	L7	L1 near25 L4		0
<input type="checkbox"/>	L6	L1 near15 L4		0
<input type="checkbox"/>	L5	L1 same L4		2
<input type="checkbox"/>	L4	((strand near3 specific) near7 \$nuclease)		583
<input type="checkbox"/>	L3	(recombinase)		4693
<input type="checkbox"/>	L2	((recombin\$)near10(recursive or recursively or repeat or repeats or repeated or repeating or iterative or iteration or iterating))		3260
<input type="checkbox"/>	L1	((sequence or domain) near10 (evolution or evolve or evolved or evolves or evolving or shuffle or shuffled or shuffles or shuffling))		5932

END OF SEARCH HISTORY

File Copy
09/08/06, 171
updated

PALM INTRANET

Day : Monday
Date: 12/6/2004
Time: 10:53:29

Inventor Name Search

Enter the first few letters of the Inventor's Last Name.

Additionally, enter the first few letters of the Inventor's First name.

Last Name

First Name

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Potential DP

- x 09/133,934 → 6,074,853 → no use of second pair for second domain evolution, or for use of ss-specific endonuclease.
- x 09/288,586 → 6,200,812 → " " " " " " " " " " .
- x 09/373,347 → ABN
- x 08/910,367 → 5,948,653 → no use of ss-specific nuclease to form a nicked target sequence; no use of second pair, second domain evolution.

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